
Release 3.1A John F. Collins, Biocomputing Research Unit.
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MSPrch_n n.a. - n.a. database search, using Smith-Waterman algorithm
Run on: Thu Jan 13 23:39:42 2000; MasPar time 73.29 Seconds
Tabular output not generated. 11/9.858 Million cell updates/sec
Title: >US-08-978-217-15
Description: (2701-3700) from US08978217.seq (4 of 10)
Perfect Score: 1000
N.A. Sequence: 2701 TCATTAAACGAGCGAGCGCT.....GGCTCCTGCTCCTCGACTA 3700
Comp: AGTAATTGGTGGTCCGGA.....CCGAGGAGGAGGAGGCTGAT

Scoring table: TABLE default
Gap 6
Nmatch STD : Dbase 0; Query 0
Searched: 165362 segs, 43234748 bases x 2
Post-processing: Minimum Match 0%
Listing first 45 summaries
Database: n-issued
1:5A_COMB 2:5B_COMB 3:5C_COMB 4:PCT9_COMB 5:Backfiles1
Statistics: Mean 8.645; Variance 5.129; scale 1.686

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length | DB | ID | Description | Pred. No. |
|------------|-------|-------------|--------|----|------------|------------------------|-----------|
| 1 | 70 | 7.0 | 1920 | 3 | US-08-746- | Sequence 1, Applicatio | 1.01e-28 |
| 2 | 49 | 4.9 | 7218 | 2 | US-08-233- | Sequence 14, Applicati | 1.98e-15 |
| 3 | 48 | 4.8 | 11236 | 1 | US-07-853- | Sequence 1, Applicatio | 8.12e-15 |
| 4 | 43 | 4.3 | 215 | 1 | US-08-238- | Sequence 5, Applicatio | 8.53e-12 |
| 5 | 42 | 4.2 | 377 | 3 | US-08-332- | Sequence 1, Applicatio | 3.36e-11 |
| 6 | 41 | 4.1 | 965 | 3 | US-08-388- | Sequence 22, Applicati | 1.32e-10 |
| 7 | 38 | 3.8 | 252 | 3 | US-08-332- | Sequence 28, Applicati | 7.51e-09 |
| 8 | 38 | 3.8 | 329 | 3 | US-08-332- | Sequence 29, Applicati | 7.51e-09 |
| 9 | 38 | 3.8 | 7218 | 2 | US-08-233- | Sequence 14, Applicati | 7.51e-09 |
| 10 | 37 | 3.7 | 2244 | 4 | PCT-US95-0 | Sequence 1, Applicatio | 2.84e-08 |
| 11 | 37 | 3.7 | 2244 | 4 | US-08-203- | Sequence 1, Applicatio | 2.84e-08 |
| 12 | 36 | 3.6 | 276 | 3 | US-08-332- | Sequence 30, Applicati | 1.06e-07 |
| 13 | 36 | 3.6 | 941 | 3 | US-08-203- | Sequence 3, Applicatio | 1.06e-07 |
| 14 | 36 | 3.6 | 941 | 4 | PCT-US95-0 | Sequence 22, Applicati | 1.06e-07 |
| 15 | 36 | 3.6 | 965 | 3 | US-08-388- | Sequence 1, Applicatio | 1.06e-07 |
| 16 | 36 | 3.6 | 3680 | 4 | PCT-US95-0 | Sequence 1, Applicatio | 1.06e-07 |
| 17 | 36 | 3.6 | 3680 | 4 | US-08-759- | Sequence 1, Applicatio | 1.06e-07 |
| 18 | 35 | 3.5 | 36 | 2 | US-08-068- | Sequence 8, Applicatio | 3.95e-07 |
| 19 | 34 | 3.4 | 54 | 4 | PCT-US92-0 | Sequence 2, Applicatio | 1.45e-06 |
| 20 | 34 | 3.4 | 183 | 4 | PCT-US95-0 | Sequence 3, Applicatio | 1.45e-06 |

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| 21 | 34 | 3.4 | 183 | 3 | US-08-255- | Sequence 3, Applicatio | 1.45e-06 |
| 22 | 34 | 3.4 | 710 | 4 | PCT-US95-0 | Sequence 2, Applicatio | 1.45e-06 |
| 23 | 34 | 3.4 | 710 | 3 | US-08-255- | Sequence 2, Applicatio | 1.45e-06 |
| 24 | 34 | 3.4 | 1505 | 1 | US-07-915- | Sequence 1, Applicatio | 1.45e-06 |
| 25 | 34 | 3.4 | 4155 | 3 | US-08-611- | Sequence 1, Applicatio | 1.45e-06 |
| 26 | 34 | 3.4 | 4155 | 1 | US-08-304- | Sequence 1, Applicatio | 1.45e-06 |
| 27 | 34 | 3.4 | 4155 | 4 | PCT-US92-0 | Sequence 1, Applicatio | 1.45e-06 |
| 28 | 34 | 3.4 | 4155 | 1 | US-08-158- | Sequence 1, Applicatio | 1.45e-06 |
| 29 | 34 | 3.4 | 4155 | 1 | US-08-316- | Sequence 1, Applicatio | 1.45e-06 |
| 30 | 34 | 3.4 | 4155 | 1 | US-07-675- | Sequence 1, Applicatio | 1.45e-06 |
| 31 | 34 | 3.4 | 4155 | 1 | US-07-675- | Sequence 1, Applicatio | 1.45e-06 |
| 32 | 34 | 3.4 | 4155 | 5 | 5281530-2 | Patent No. 5281530. | 1.45e-06 |
| 33 | 34 | 3.4 | 4155 | 5 | 5426049-2 | Patent No. 5426049. | 1.45e-06 |
| 34 | 34 | 3.4 | 4155 | 1 | US-08-053- | Sequence 1, Applicatio | 1.45e-06 |
| 35 | 33 | 3.3 | 215 | 1 | US-08-238- | Sequence 5, Applicatio | 5.26e-06 |
| 36 | 33 | 3.3 | 297 | 1 | US-08-222- | Sequence 33, Applicati | 5.26e-06 |
| 37 | 33 | 3.3 | 3596 | 3 | US-08-779- | Sequence 5, Applicatio | 5.26e-06 |
| 38 | 33 | 3.3 | 3632 | 3 | US-08-779- | Sequence 5, Applicatio | 5.26e-06 |
| 39 | 33 | 3.3 | 3632 | 3 | PCT-US93-0 | Sequence 4, Applicatio | 5.26e-06 |
| 40 | 33 | 3.3 | 10747 | 4 | PCT-US93-0 | Sequence 1, Applicatio | 5.26e-06 |
| 41 | 33 | 3.3 | 24979 | 4 | PCT-US93-0 | Sequence 3, Applicatio | 5.26e-06 |
| 42 | 32 | 3.2 | 38 | 2 | PCT-US95-0 | Sequence 1, Applicatio | 1.89e-05 |
| 43 | 32 | 3.2 | 38 | 2 | US-08-563- | Sequence 1, Applicatio | 1.89e-05 |
| 44 | 32 | 3.2 | 605 | 4 | PCT-US96-0 | Sequence 89, Applicati | 1.89e-05 |
| 45 | 32 | 3.2 | 3535 | 3 | US-08-618- | Sequence 1, Applicatio | 1.89e-05 |

ALIGNMENTS

RESULT 1
ID US-08-746-789A-1 STANDARD: DNA; UNC: 1920 BP.
AC xxxxxx
DE Sequence 1, Application US/08746789A
CC Sequence 1, Application US/08746789A
CC Patent No. 5789200
CC GENERAL INFORMATION:
CC APPLICANT: Ismail Kola, Martin J. Tyms, Christine DeBuck
CC TITLE OF INVENTION: A No. 5789200e1 Human ETS Family Member, ELF3
CC NUMBER OF SEQUENCES: 4
CC CORRESPONDENCE ADDRESS:
CC ADDRESSEE: SmithKline Beecham Corporation
CC STREET: 709 Swedeland Road, P.O. Box 1539
CC CITY: King of Prussia
CC STATE: PA
CC COUNTRY: USA
CC ZIP: 19406-0939
CC COMPUTER READABLE FORM:
CC MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CC COMPUTER: IBM 486
CC OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
CC SOFTWARE: MICROSOFT WORD
CC CURRENT APPLICATION DATA:
CC APPLICATION NUMBER: US/08/746,789A
CC FILING DATE: No. 5789200e1 15, 1996
CC CLASSIFICATION: 514
CC PRIOR APPLICATION DATA:
CC APPLICATION NUMBER:
CC FILING DATE:
CC ATTORNEY/AGENT INFORMATION:
CC NAME: William T. Han
CC REGISTRATION NUMBER: 34,344
CC REFERENCE/DOCKET INFORMATION: ATG 50024
CC TELECOMMUNICATION INFORMATION:
CC TELEPHONE: 610 270 5219
CC TELEFAX: 610 270 4026
CC INFORMATION FOR SEQ ID NO: 1:
CC SEQUENCE CHARACTERISTICS:
CC LENGTH: 1920
CC TYPE: Nucleic Acid
CC STRANDEDNESS: Single
CC TOPOLOGY: Linear
CC ANTI-SENSE: NO

CC MEDIUM TYPE: Floppy disk
CC COMPUTER: IBM PC compatible
CC OPERATING SYSTEM: PC-DOS/MS-DOS
CC SOFTWARE: Patentin Release #1.0, Version #1.25
CC CURRENT APPLICATION DATA:
CC APPLICATION NUMBER: US/08/203,532F
CC FILING DATE:
CC CLASSIFICATION: 435
CC ATTORNEY/AGENT INFORMATION:
CC NAME: Goltick, Mary E.
CC REGISTRATION NUMBER: 34829
CC REFERENCE/DOCKET NUMBER: 22311/00114
CC TELECOMMUNICATION INFORMATION:
CC TELEPHONE: (216) 622-8200
CC TELEFAX: (216) 241-0816
CC
CC INFORMATION FOR SEQ ID NO: 3:
CC SEQUENCE CHARACTERISTICS:
CC LENGTH: 941 base pairs
CC TYPE: nucleic acid
CC STRANDEDNESS: both
CC TOPOLOGY: linear
CC MOLECULE TYPE: CDNA
CC HYPOTHETICAL: NO
CC ANTI-SENSE: NO
CC FEATURE:
CC NAME/KEY: CDS
CC LOCATION: 33..941
CC
SQ SEQUENCE 941 BP; 252 A; 293 C; 242 G; 154 T; 0 OTHER.

Query Match 3.6%; Score 36; DB 3; Length 941;
Best Local Similarity 89.1%; Pred. No. 1.06e-07;
Matches 41; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

Db 234 CACCACCACCACCACCACCATCACCACCATCAGCAGCAGC 279
|||
Cp 3430 CACCACCACCACCACCACCATCACCACCATCAGCAGCAGC 3385

RESULT 14
ID PCT-US95-01882A-3 STANDARD; DNA; UNC; 941 BP.
AC xxxxxx
DE Sequence 3, Application PC/TUS9501882A
CC Sequence 3, Application PC/TUS9501882A
CC GENERAL INFORMATION:
CC APPLICANT: Gorski, David H.
CC APPLICANT: Walsh, Kenneth
CC TITLE OF INVENTION: Growth Arrest Homeobox Gene
CC NUMBER OF SEQUENCES: 4
CC CORRESPONDENCE ADDRESS:
CC ADDRESSEE: Calfee, Halter, and Griswold
CC STREET: 800 Superior Avenue
CC City: Cleveland
CC STATE: Ohio
CC COUNTRY: U.S.A.
CC ZIP: 44114-2688
CC COMPUTER READABLE FORM:
CC MEDIUM TYPE: Floppy disk
CC COMPUTER: IBM PC compatible
CC OPERATING SYSTEM: PC-DOS/MS-DOS
CC SOFTWARE: Patentin Release #1.0, Version #1.25
CC CURRENT APPLICATION DATA:
CC APPLICATION NUMBER: PCT/US95/01882A
CC FILING DATE:
CC CLASSIFICATION:
CC ATTORNEY/AGENT INFORMATION:
CC NAME: Goltick, Mary E.
CC REGISTRATION NUMBER: 34829
CC REFERENCE/DOCKET NUMBER: 22311/00114
CC TELECOMMUNICATION INFORMATION:
CC TELEPHONE: (216) 622-8200
CC TELEFAX: (216) 241-0816
CC

CC TELEX: 980499
CC INFORMATION FOR SEQ ID NO: 3:
CC SEQUENCE CHARACTERISTICS:
CC LENGTH: 941 base pairs
CC TYPE: nucleic acid
CC STRANDEDNESS: both
CC TOPOLOGY: linear
CC MOLECULE TYPE: CDNA
CC HYPOTHETICAL: NO
CC ANTI-SENSE: NO
CC FEATURE:
CC NAME/KEY: CDS
CC LOCATION: 33..941
CC
SQ SEQUENCE 941 BP; 252 A; 293 C; 242 G; 154 T; 0 OTHER.

Query Match 3.6%; Score 36; DB 4; Length 941;
Best Local Similarity 89.1%; Pred. No. 1.06e-07;
Matches 41; Conservative 0; Mismatches 5; Indels 0; Gaps 0;

Db 234 CACCACCACCACCACCACCATCACCACCATCAGCAGCAGC 279
|||
Cp 3430 CACCACCACCACCACCACCATCACCACCATCAGCAGCAGC 3385

RESULT 15
ID US-08-388-672A-22 STANDARD; DNA; UNC; 965 BP.
AC xxxxxx
DE Sequence 22, Application US/08388672A
CC Sequence 22, Application US/08388672A
CC Patent No. 5795961
CC GENERAL INFORMATION:
CC APPLICANT: Wallace, T. Paul
CC APPLICANT: Harris, William J.
CC APPLICANT: Carr, Frank J.
CC APPLICANT: Old, Lloyd J.
CC APPLICANT: Welt, Sydney
CC APPLICANT: Kitamura, Kunio
CC TITLE OF INVENTION: Recombinant Human Anti-Lewis B
CC NUMBER OF SEQUENCES: 25
CC CORRESPONDENCE ADDRESS:
CC ADDRESSEE: Felle and Lynch
CC STREET: 805 Third Avenue
CC City: New York
CC STATE: New York
CC COUNTRY: U.S.A.
CC ZIP: 10022
CC COMPUTER READABLE FORM:
CC MEDIUM TYPE: Floppy disk
CC COMPUTER: IBM PC compatible
CC OPERATING SYSTEM: PC-DOS/MS-DOS
CC SOFTWARE: Patentin Release #1.0, Version #1.30
CC CURRENT APPLICATION DATA:
CC APPLICATION NUMBER: US/08/388,672A
CC FILING DATE: 14-FEB-1995
CC CLASSIFICATION:
CC ATTORNEY/AGENT INFORMATION:
CC NAME: Hanson, No. 5795961man D.
CC REGISTRATION NUMBER: 30,946
CC REFERENCE/DOCKET NUMBER: LUD 5409
CC TELECOMMUNICATION INFORMATION:
CC TELEPHONE: 212-688-9200
CC TELEFAX: 212-838-3884
CC INFORMATION FOR SEQ ID NO: 22:
CC SEQUENCE CHARACTERISTICS:
CC LENGTH: 965 base pairs
CC TYPE: nucleic acid
CC STRANDEDNESS: unknown
CC TOPOLOGY: unknown
CC MOLECULE TYPE: DNA (genomic)
SQ SEQUENCE 965 BP; 192 A; 170 C; 226 G; 200 T; 177 OTHER.

